## SEQUENCE LISTING



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<110> Kutyavin, Igor V.
      Milesi, David
      Hoekstra, Merl
      Epoch Biosciences, Inc.
<120> Abasic Site Endonuclease Assay
<130> 17682A-007910US
<140> US 10/645,353
<141> 2003-08-20
<150> US 60/405,642
<151> 2002-08-21
<160> 17
<170> PatentIn Ver. 2.1
<210> 1
<211> 47
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence:hairpin
      substrate structure simulating probe-target
      nucleic acid-enhancer complex
<220>
<221> modified base
<222> (47)
<223> n = c modified by 3' tail
      2-(6-oxy-2-{2-[5-hydroxy-8-(oxy-methoxy-phosphoryloxy)-
      octylcarbamoyl]-ethyl}-3-oxo-3H-xanthen-9-yl)-benzoate
      (structure #2)
<400> 1
                                                                    47
gccacattgg aagccaatgt ggcgggcaag gaccgaaggt ccttgcn
<210> 2
<211> 5
<212> PRT
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<223> Description of Artificial Sequence:flexible
      polypeptide chain probe-enhancer linker
<400> 2
Gly Ser Ser Ser Ser
  1
<210> 3
<211> 24
<212> DNA
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<213> Artificial Sequence

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<220'>
<223> Description of Artificial Sequence:target
     oligonucleotide
<400> 3
                                                                   24
aatgtggcgg gcaaggaccg agtc
<210> 4
<211> 11
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence:11-mer probe
      complementary to target oligonucleotide
<220>
<221> modified base
<222> (1)
<223> n = a modified by 5' conjugated quencher
     phosphoric acid 1-(4-{[4-(2-chloro-4-nitro-phenylazo)-
     phenyl]-methyl-amino}-butyryl)-5-hydroxymethyl-pyrrolidin-
      3-yl ester methyl ester (structure #15)
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<221> modified base
<222> (11)
<223> n = t modified by 3' tail of fluorescein (FAM) and linker 4,5-
     dichloro-2-(4,7-dichloro-6-oxy-2-{3-[4-hydroxy-2-(oxy-methoxy-
     phosphoryloxymethyl)-pyrrolidin-1-yl]-3-oxo-propyl}-5-methyl-3-
     oxo-3H-xanthen-9-yl)-benzoate (structure #8)
<400> 4
                                                                   11
nctcggtcct n
<210> 5
<211> 11
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence:enhancer
      oligonucleotide for 1 base gap between target
     oligonucleotide and duplexes of probe and enhancer
<400> 5
                                                                   11
cccgccacat t
<210> 6
<211> 11
<212> DNA
<213> Artificial Sequence
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      oligonucleotide for 0 base gap between target
      oligonucleotide and duplexes of probe and enhancer
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<400> 6
                                                                    11
gcccgccaca t
<210> 7
<211> 10
<212> DNA
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<220>
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      oligonucleotide for 2 base gap between target
      oligonucleotide and duplexes of probe and enhancer
<400> 7
                                                                    10
ccgccacatt
<210> 8
<211> 41
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:41-mer fully
      matched DNA target sequence
<400> 8
                                                                    41
agtcacagtc ggtgccaatg tggcgggcaa ggaccgagtc g
<210> 9
<211> 22
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence:enhancer
      oligonucleotide
<400> 9
                                                                    22
gccacattgg caccgactgt ga
<210> 10
<211> 14
<212> DNA
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<223> Description of Artificial Sequence:14-mer
      oligonucleotide probe
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<221> modified_base
<222> (1)
<223> n = a modified by 5' conjugated quencher
      phosphoric acid 1-(4-{[4-(2-chloro-4-nitro-phenylazo)-
      phenyl]-methyl-amino}-butyryl)-5-hydroxymethyl-pyrrolidin-
      3-yl ester methyl ester (structure #15)
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<220">
<221> modified_base
<222> (14)
<223> n = t modified by 3' tail of fluorescein (FAM) and linker 4,5-
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     phosphoryloxymethyl)-pyrrolidin-1-yl]-3-oxo-propyl}-5-methyl-3-
     oxo-3H-xanthen-9-yl)-benzoate (structure #8)
<400> 10
nctcggtcct tgcn
                                                                   14
<210> 11
<211> 10
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence:10-mer
     oligonucleotide probe
<220>
<221> modified base
<222> (1)
<223> n = a modified by 5' conjugated quencher
     phosphoric acid 1-(4-{[4-(2-chloro-4-nitro-phenylazo)-
     phenyl]-methyl-amino}-butyryl)-5-hydroxymethyl-pyrrolidin-
     3-yl ester methyl ester (structure #15)
<220>
<221> modified base
<222> (10)
<223> n = t modified by 3' tail of fluorescein (FAM) and linker 4,5-
     dichloro-2-(4,7-dichloro-6-oxy-2-{3-[4-hydroxy-2-(oxy-methoxy-
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     oxo-3H-xanthen-9-yl)-benzoate (structure #8)
<400> 11
                                                                   10
ngtccttgcn
<210> 12
<211> 27
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: fragment of
      target sequence around single nucleotide
     polymorphism in human genomic DNA
<400> 12
                                                                    27
aaagagacac ggacayatca atccatc
<210> 13
<211> 20
<212> DNA
<213> Artificial Sequence
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<220">
 <223> Description of Artificial Sequence:asymmetric PCR
      amplification forward primer
 <400> 13
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 caaactttgt ccttggtcta
 <210> 14
 <211> 20
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence:asymmetric PCR
      amplification reverse primer
 <400> 14
                                                                     20
 ttcttttacc actccccctt
 <210> 15
 <211> 13
 <212> DNA
 <213> Artificial Sequence
 <223> Description of Artificial Sequence:complementary
       target oligodeoxyribonucleotide (ODN)
 <400> 15
                                                                     13
 caaggaccga gtc
 <210> 16
 <211> 11
 <212> DNA
 <213> Artificial Sequence
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 <223> Description of Artificial
      Sequence:oligodeoxyribonucleotide (ODN) probe
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 <221> modified_base
 <222> (1)
 <223> n = a modified by 5' conjugated quencher
      phosphoric acid 1-(4-{[4-(2-chloro-4-nitro-phenylazo)-
      phenyl] -methyl -amino} -butyryl) -5-hydroxymethyl -pyrrolidin-
      3-yl ester methyl ester (structure #15)
<220>
 <221> modified base
 <222> (11)
 <223> n = t modified by 3' tail of fluorescein (FAM) and linker 4,5-
      dichloro-2-(4,7-dichloro-6-oxy-2-{3-[4-hydroxy-2-(oxy-methoxy-
      phosphoryloxymethyl)-pyrrolidin-1-yl]-3-oxo-propyl}-5-methyl-3-
      oxo-3H-xanthen-9-yl)-benzoate (structure #8)
 <400> 16
                                                                     11
nctcggtcct n
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<210 > 17
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<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial
     Sequence:oligodeoxyribonucleotide (ODN) probe
<220>
<221> modified_base
<222> (1)
<223> n = a modified by 5' conjugated quencher
     phosphoric acid 1-(4-{[4-(2-chloro-4-nitro-phenylazo)-
     phenyl]-methyl-amino}-butyryl)-5-hydroxymethyl-pyrrolidin-
     3-yl ester methyl ester (structure #15)
<400> 17
nctcggtcct t
                                                                   11
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